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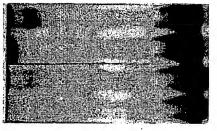
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### FIGURE 1

nt14-58+MBP-Rep78(.5) nt57-106+MBP-Rep78(.5)



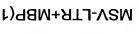
(f.)87q9A-9AH+80f-4lin (1)98M+801-4tJn (f.)87q9A-9AM+61-1487in nt7841-13+MBP(1)

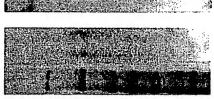




nt7841-106+MBP-Rep78(1) nt7841-106+MBP-Rep78(.1) (1)98M+301-1487in MSV-LTR+MBP-Rep78(1) MSV-LTR+MBP(1)



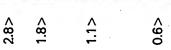


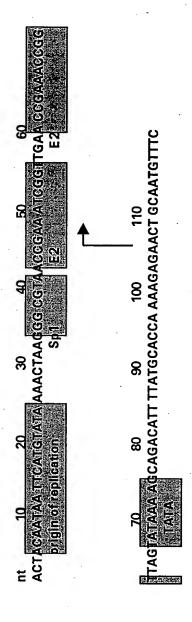


TR+MBP-Rep78(1)

Rep78 selected fragments

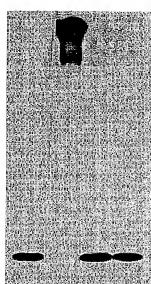
IH ms8,1 is 7 X 81-V9H

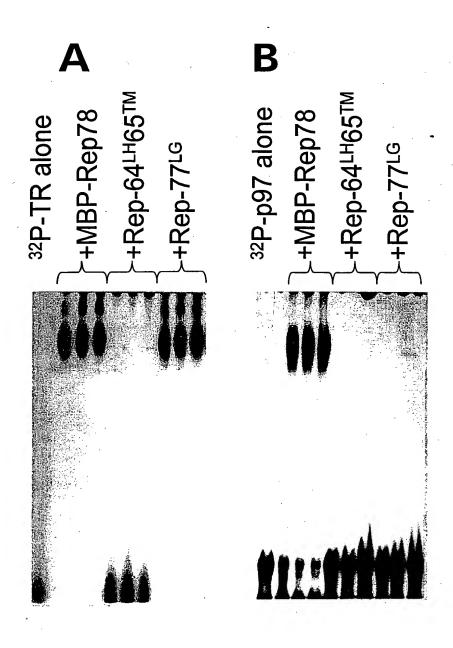


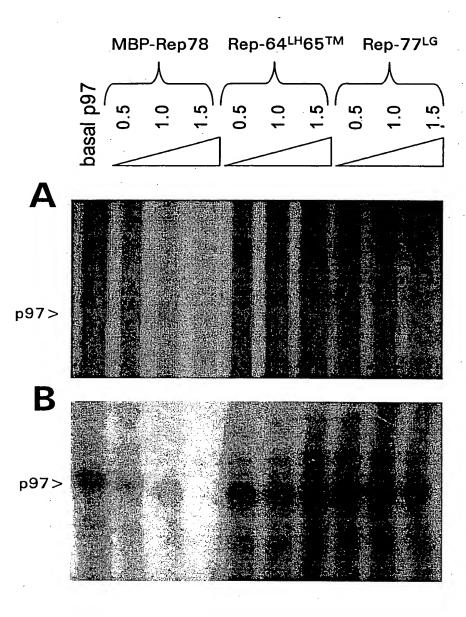


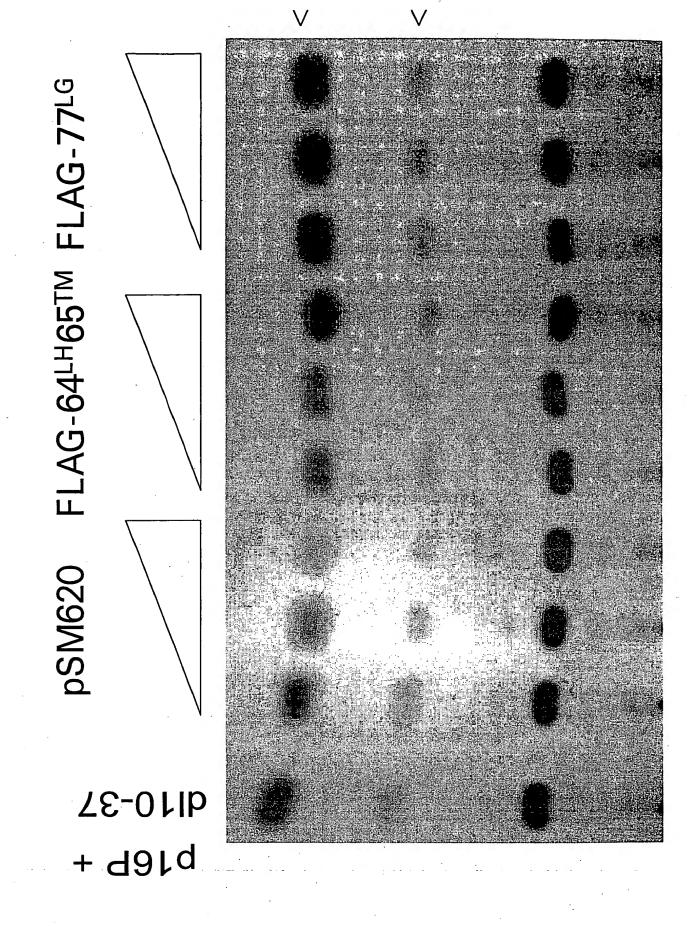
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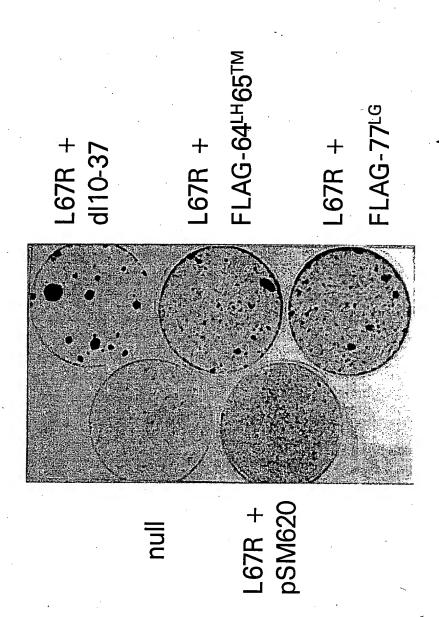
strand +MBP-Rep78strand + MBPstrand +MBP-Rep78

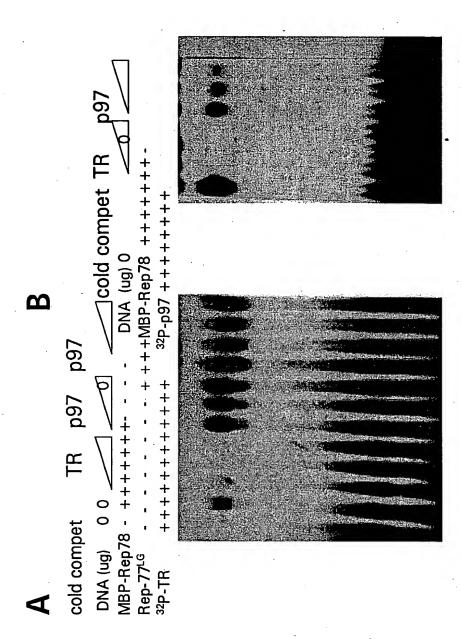


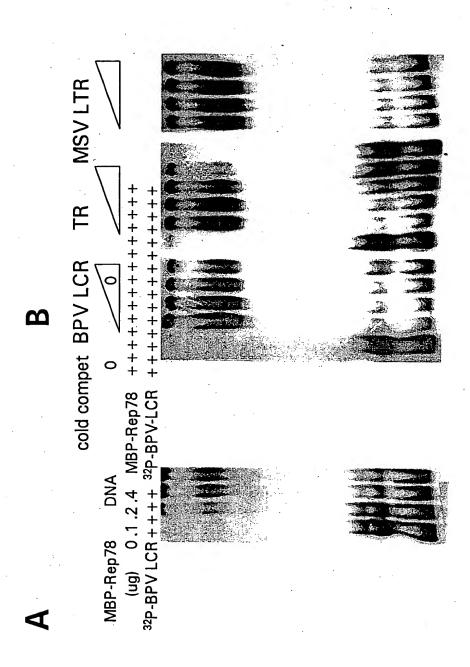


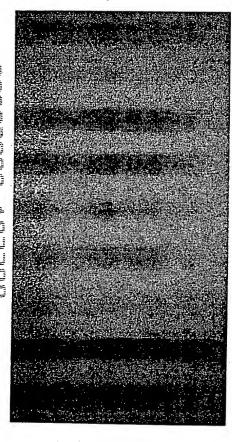












MBP

MBP-Rep78

Rep-64<sup>LH</sup>65<sup>TM</sup>

Rep-77<sup>LG</sup>

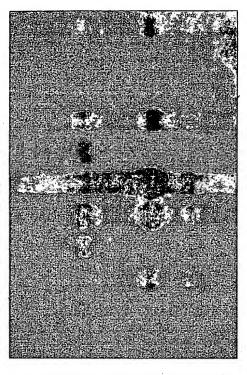
Rep-79<sup>FA</sup>

Rep-105<sup>LS</sup>

Rep-192<sup>HG</sup>

Rep-331WG

Rep-340<sup>KA</sup>



marker blank Ad alone

pSM620/Sph

FLAG-64<sup>LH</sup>65<sup>TM</sup>

FLAG-77<sup>LG</sup>

FLAG-79FA

FLAG-105<sup>LS</sup>

FLAG-192HG

FLAG-331WG

FLAG340<sup>KA</sup>

<sup>32</sup>P-TR DNA

MBP

MBP-Rep78

Rep-64<sup>LH</sup>65<sup>™</sup>

Rep-77<sup>LG</sup>

Rep-79<sup>FA</sup>

Rep-105<sup>LS</sup>

Rep-192<sup>HG</sup>

Rep-331WG

Rep-340<sup>KA</sup>

# Figure 13

# Figure 14

oncogenic foci
plasmids transfected

34	<b>∞</b>	19	25	4
pL67R + 334 (no Rep)	pL67R + 620/Sph (wild type)	$pL67R + Rep77^{LG}$	$pL67R + Rep79^{FA}$	$pL67R + Rep192^{HG}$

#### FIGURE 15A

aaaggtcgcc gcgacaccat gtccccagcg gccgagaagg gcaccctga agtaaggccc atgcacgtgc cagattcgcg tggttcgcgg tgctacatcc aatatggaac atctccttca ggaaagatta ctcgtggaca gaggacattt tatgcggctt tggctgtttg gacaagatgg aactcaacga aaagccattc atagacccga acccgccgtc gtgcccttct agaggagt tacgtcata tgaagcggg tctgatgcg cagcatctg gagcgcgcag gtctccattt ccgggcgacc tgacgtgaat gcgacatttt tgtgattaag gaactgggtg tttgccaaac gattgagcag gcgccgtgtg ctacttccac tttcctgagt ggtggatgag ggcgtggact gttggtggcg gaatcccaat ggtcgggtgg ggcctcatac ggacaatgcg gaacaccatc gcagcccgtg cgatccccaa agcccacact cgactgtgtc ggagtcggcc ctcggcccag gattgacggg atttgaactc ctcactgagg gtgagcgagc ggtggagtcg tgagtgtttt agcacgcagg tttacgagat acagctttgt tgacggaatg tttgggacg tcgagccgac atctgaatct agggagagag ggaacaaggt agctccagtg agcgtaaacg aagagaatca acatggagct aggaggacca aggctgcctt tggtgggcca taaacgggta toggcaagag cggaggccat ttcccttcaa ccaaggtcgt aatgcaagtc tgtgcgccgt ggatgttcaa cgctcgctcg ggcggcctca ttcctggagg agaggtcacg agcccgagtg atgccggggt ggcatttctg cgcgactttc tctgacatgg caatttgaga taccgcggga aaatccatgg gccggaggcg acccagcctg aatctcacgg gagcagaaca cagtggatcc tcccaaatca cccgactacc tcagccaggt attttggaac acgaaaagt accaacatcg aatgagaact aagatgaccg gtggaccaga aacaccaaca ttgcaagacc cctctctgcg tgggtattta gatttataaa gctttgcccg tcactagggg greetgtatt cgcagccgcc gcatctgccc gccgccagat gaagctgcag gctcccaaa tttctttgtg tcagagaätt cgcctgtttg caccggggtg cagaaatggc gcagacgcag atcaaaact ctcggagaag caactcgcgg taaaaccgcc gggatgggcc taccgggaag aaactggacc 99a99a9999 caaggtgcgc cgtcacctcc ccagcagccg ttggccactc cgacgcccgg gccaactcca ggttagggag gtggtcacgc ggtttgaacg accttgacgg aatgggagtt ccgtggccga cggaggccct tcgtggaaac aaaaactgat tcacaaagac agtatttaag cgcacgtgtc aggggattac tgagcctgac ccaattactt cggtgatcag atgeggeete ccagcaatcg ggcctgcaac tgatctggtg ccgtctttct acgggtgcgt tcggaggaag ctcccgtgat ccttcgaaca 181 241 301 361 481 541 421 601 661 721 781 841 901 961 1021 1081 1141 1201 1261 1321 1381 1441

#### FIGURE 15B

att D gttct agaggccggt gccagcagcc ctgacccca U D tgctttcccg g caaaacaaat tgctacattc aatgtggatt cgatggttat gcg tcagttgcg gagagaatg tggctacag attcctccg a g gccaagaaa cggtgggca gccggtca gcaccga tcacgcag gggtctt aatcagg acttcaa actgggg ggagtca gaagct tcgaca gcctta agagg ggatgcctcc tcagaaactg cgatctggtc gtatggctgc gacagtggtg gaccggcagc tttcaggagc caggcgaaaa ccgggaaaa ggaaaggcgg actaatacga ggagtgggta atcaccacca attccagcc gggtattttg atcaacaaca gtcaaagagg gttcaggtgt ctgaacaacg agactttttc aaagggtgga agacaggtac cagacaatgc ctgtttagag D ggtgcgcgag gactcagta acgacagca acaagggag taacattcaa ggcgcatcaa atacctcacc agcagtcttc ctcgggaacc tggtctggga gggcgccgac cgacagagtc ctacaaacaa cacccttgg gcaaagactc taccagcacg tgttccctg aaaaggcgta b tttaaatcag U cgacgcggag taagacggct tggagacgca ಡ agcccaaacg tcaactacgc ಡ ಥ δ aacggactcg cggcataag caaagccta aggaagtca tctacgtca gacagaaag gcactgcct gaaggaata cacagtatgg tcctcggctc gtcaccaagc aatctgatgc caataaatga gcccgcagag cggacccttc cgagcacgta acaaccacgc acctcggacg aggaacctgt cagactcctc ttggtcagac cagcccctc acaataacga catggatggg acaaccacct ttggctacag cacgtgactg tcaagctctt ccaataacct gagcatgaat gaagcttcga ttcactcacg tctgtcgtca ccagacgctt cactctctct gatataagtg ctcccgtacg ttcatggtgc tgcgattcca aatcactact cácttttcac agactcaact acgacgattg ctttgggaag ggttgaggtg cagtgacgca aaatatctgc tcaacccgtt gggaaaggtg catctitgaa ggctcgagga caccaccaaa acaagtacct ccgcggccct tacctcaagt tttgggggca ggcctggttg cctgtggagc agattgaatt cagccaccag ccaatiggcag cccacctaca gtcagacgcg cgtgggcatg gcagacgtc ggagtaccag agatacgtct ctcgaacgac attccactgc ccgacccaag tgacggtacg tggatcatga accegeeee gccatcgac atcagaattc gtcagaatc atcatatcat ggatgactg cctggcccac cttcctgggt gaggcagacg agacaacccg tgaacctctg agagcactct tgcaagaaa gcctctcgga cagtggcgca aaattggcat ctgggccctg aggatcacgt gttctcgtca cttccagatt 2581 2401 2881 2941 3001 121 3181 241 301 2461 2521 2641 2701 2761 2821 3061 1741 921 981 041 101 2221 2281 2341 1801 1861 2161

### FIGURE 15C

aaggctcaga gctacgctca attacttgag ctcaggccgg accgccagca gccacaagga tctggcagga teggaacaac gaggcaacag agtttgcttc gagctgcaga aacaagtctg agattctcat cccattggca taattcgttt tacgtagata ttggccactc cgacgcccgg acggacattt ctggagcta tgcgtaccg gccaa tctcagatgc ttccacagca cagtacctgt cttcagttt ggaccctgtt tactcgtgga gccatggcaa tttgggaagc gaagaggaaa aacctccaga ccaggcatgg ccacacacgg agtgcggcaa gatcgagtgg agagcctcgc cctcctccac ttccaactac tttccatggc agtgatggag aaaggtcgcc ataaaccgtt agaggagtg ggacgttcct ggttctcatc tctcatcgac gcagtcaagg gtactttcct caacagtgaa gattacagac ggtgaatccg tgtatctacc tcagcgtgga ttcagtacac ctggcttcct aggcgttctt ggcaaagatt acttaaacac gaccaccttc gcgtgtattc ttgttaatca tcttatctag aggaaccct ccgggcgacc gagcgcgcag acacttttga actgcctgga tcatgaatcc gaaccaccac agtctaggaa cggataacaa gagactctct ctcagagcgg aaaaggtcat agtatggttc tcaacaca ggcccatctg cgaatccttc cgggacacgg aatcccgaaa gatactaatg ctgtaattgc cgtatttctt attaactaca ctcactgagg gtggattcgg gtgagcgagc accttcagct ctggaccgtc actccaagtg attcgggacc ctcaatggca tcttcatttt aagacatctg aagtttttc gtgaacattg gctacggagc accgcagatg taccttcagg cagtactcca cgggttaatc cgctcgctcg ccctcatgg ccggtacctg caaacgctgg acttaccgtg gactcgtaat ttggtctctg gġcggcctca agtaggacgc aaacaacttt cagaacaac agcgagtgac caagtaccac cgatgaagaa ttaatcgtgg cctctctgcg cagccagagt gcgagtatca gaaaacaaat caatcccgtg acaagcagct cagagatgtg tcacccctct caagaacacc cttcatcaca aggaaaacag ccagatacct cagttgaact agtagcatgg gctttgcccg 3421 3481 4561 3361 3541 3601 3661 3721 3781 3841 3901 4021 4081 4141 4321 4381 4441 4501 4621 3961 4201 4261

### FIGURE 16

MPGFYEIVIKVPSDLDGHLPGISDSFVNWVAEKEWELPPDSDMD LNLIEQAPLTVAEKLQRDFLTEWRRVSKAPEALFFVQFEKGESYFHMHVLVETTGVKS MVLGRFLSQIREKLIQRIYRGIEPTLPNWFAVTKTRNGAGGGNKVVDECYIPNYLLPK TQPELQWAWTNMEQYLSACLNLTERKRLVAQHLTHVSQTQEQNKENQNPNSDAPVIRS **TKTÄPDYLVGQQPVEDISSNRIYKILELNGYDPQYAASVFLGWATKKFGKRNTIWLFG** PATTGKTNIAEAIAHTVPFYGCVNWTNENFPFNDCVDKMVIWWEEGKMTAKVVESAKA TRRLDHDFGKVTKQEVKDFFRWAKDHVVEVEHEFYVKKGGAKKRPAPSDADISEPKRV **RESVAQPSTSDAEASINYADRYQNKCSRHVGMNLMLFPCRQCERMNQNSNICFTHGQK** KTSARYMELVGWLVDKGITSEKQWIQEDQASYISFNAASNSRSQIKAALDNAGKIMSL ILGGSKVRVDQKCKSSAQIDPTPVIVTSNTNMCAVIDGNSTTFEHQQPLQDRMFKFEL DCLECFPVSESQPVSVVKKAYQKLCYIHHIMGKVPDACTACDLVNVDLDDCIFEQ